

EVANS ET AL.
USSN: 08/244,857
Filed: June 14, 1994
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No fee is deemed necessary in connection with the filing of this Preliminary Amendment. However, if any fee is required, authorization is given to charge the amount of this fee to Deposit Account No. 16-2460.

Respectfully submitted,

10/12/94
Date

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PRETTY, SCHROEDER,
BRUEGGEMANN & CLARK
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SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Evans, Ronald M.
Mangelsdorf, David J.
Heyman, Richard A.
Boehm, Marcus F.
Eichele, Gregor
Thaller, Christina
- (ii) TITLE OF INVENTION: MEANS FOR THE MODULATION OF PROCESSES
MEDIATED BY RETINOID RECEPTORS AND COMPOUNDS USEFUL
THEREFOR
- (iii) NUMBER OF SEQUENCES: 2
- (iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: Pretty, Schroeder, Brueggemann & Clark
(B) STREET: 444 South Flower Street, Suite 2000
(C) CITY: Los Angeles
(D) STATE: CA
(E) COUNTRY: USA
(F) ZIP: 90071
- (v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- (vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER: US 08/244,857
(B) FILING DATE: 14-JUN-1994
(C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
(A) APPLICATION NUMBER: US 07/809,980
(B) FILING DATE: 18-DEC-1991
- (vii) PRIOR APPLICATION DATA:
(A) APPLICATION NUMBER: WO 93/11755
(B) FILING DATE: 18-DEC-1992
- (viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: Reiter, Stephen E.
(B) REGISTRATION NUMBER: 31,192
(C) REFERENCE/DOCKET NUMBER: P41 9755
- (ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: 619-546-4737
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(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 71 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: unknown
- (ii) MOLECULE TYPE: protein
- (v) FRAGMENT TYPE: internal

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Cys	Xaa	Xaa	Cys	Xaa	Xaa	Asp	Xaa	Ala	Xaa	Gly	Xaa	Tyr	Xaa	Xaa	Xaa
1				5					10					15	
Xaa	Cys	Xaa	Xaa	Cys	Lys	Xaa	Phe	Phe	Xaa	Arg	Xaa	Xaa	Xaa	Xaa	Xaa
			20					25					30		
Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Cys	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Xaa	Cys
			35				40					45			
Xaa	Xaa	Xaa	Lys	Xaa	Xaa	Arg	Xaa	Xaa	Cys	Xaa	Xaa	Cys	Arg	Xaa	Xaa
			50			55						60			
Lys	Cys	Xaa	Xaa	Xaa	Gly	Met									
65					70										

51 (2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 16 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: both
 (D) TOPOLOGY: both

(ii) MOLECULE TYPE: DNA (genomic)

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TCAGGTCATG ACCTGA